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## SEQUENCE LISTING

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<120> DNA ENCODING CANINE VON WILLEBRAND FACTOR AND METHODS OF USE

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<170> PatentIn Ver. 2.0

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Leu Leu Ile Ile Asn Ile Phe His Asp Asn Ile Asp Asp Ile Asp Ile Asp  
755 760 765 770

Asp Ile Met Val Ile Ile Val Tyr Ile Ala Asp Asn Ile Asp Ala Ile  
775 780 785

Gly Leu Ile Cys Ala Lys Thr Cys Glu Asn Tyr Asp Leu Gln Cys Met  
790 795 800

Asp Ile Ile Asp Ile Val Ile Val Tyr Ile Ala Asp Asp Ile Val Asp  
810 815 820 825

Cys Val Cys Arg Asp Arg Lys Trp Thr Cys Thr Asp His Val Cys Asp  
850 865 860

Ala Thr Cys Ser Ala Ile Gly Met Ala His Tyr Leu Thr Phe Asp Gly  
865 870 875 880

Leu Lys Tyr Leu Phe Pro Gly Glu Cys Gln Tyr Val Leu Val Gln Asp  
885 890 895

Tyr Cys Gly Ser Asn Pro Gly Thr Leu Arg Ile Leu Val Gly Asn Glu  
900 905 910

Gly Cys Ser Tyr Pro Ser Val Lys Cys Lys Lys Arg Val Thr Ile Leu  
915 920 925

Val Glu Gly Glu Ile Glu Leu Phe Asp Gly Glu Val Asn Val Lys  
930 935 940

Lys Pro Met Lys Asp Glu Thr His Phe Glu Val Val Glu Ser Gly Gln  
945 950 955 960

Tyr Val Ile Leu Leu Gly Lys Ala Leu Ser Val Val Trp Asp His  
965 970 975

Arg Leu Ser Ile Ser Val Thr Leu Lys Arg Thr Tyr Gln Glu Gln Val  
980 985 990

Cys Gly Leu Cys Gly Asn Phe Asp Gly Ile Gln Asn Asn Asp Phe Thr  
995 1000 1005

Ser Ser Ser Leu Gln Ile Glu Glu Asp Pro Val Asp Phe Gly Asn Ser  
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Trp Lys Val Asn Pro Gln Cys Ala Asp Thr Lys Lys Val Pro Leu Asp  
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Ser Ser Pro Ala Val Cys His Asn Asn Ile Met Lys Gln Thr Met Val  
1045 1050 1055

Asp Asn Ser Tyr Asp Ile Ile Thr Asp Asp Ile Ile Asp Tyr Asp  
1060 1065 1070 1075

Asp Leu Val Asp Ile Gln Ile Thr Ile Asp Ile Val Ile Tyr Asp Thr  
1075 1080 1085

Tyr Ser Cys Glu Ser Ile Gly Asp Tyr Thr Cys Ile Cys Asp Thr Ile  
1090 1095 1100

Ala Ala Tyr Ala His Val Tyr Ala Ile His Gly Ile Val Val Asp Ile

Cys Pro Ile Thr Cys Gln His Pro Glu Pro Leu Ala Cys Pro Val Gln  
1155 1160 1165

Cys Val Glu Gly Cys His Ala His Cys Pro Pro Gly Lys Ile Leu Asp  
1170 1175 1180

Glu Leu Leu Gln Thr Cys Ile Asp Pro Glu Asp Cys Pro Val Cys Glu  
1185 1190 1195 1200

Val Ala Gly Arg Arg Leu Ala Pro Gly Lys Lys Ile Ile Leu Asn Pro  
1205 1210 1215

Ser Asp Pro Glu His Cys Gln Ile Cys Asn Cys Asp Gly Val Asn Phe  
1220 1225 1230

Thr Cys Lys Ala Cys Arg Glu Pro Gly Ser Val Val Val Pro Pro Thr  
1235 1240 1245

Asp Gly Pro Ile Gly Ser Thr Thr Ser Tyr Val Glu Asp Thr Ser Glu  
1250 1255 1260

Pro Pro Leu His Asp Phe His Cys Ser Arg Leu Leu Asp Leu Val Phe  
1265 1270 1275 1280

Leu Leu Asp Gly Ser Ser Lys Leu Ser Glu Asp Glu Phe Glu Val Leu  
1285 1290 1295

Lys Val Phe Val Val Gly Met Met Glu His Leu His Ile Ser Gln Lys  
1300 1305 1310

Arg Ile Arg Val Ala Val Val Glu Tyr His Asp Gly Ser His Ala Tyr  
1315 1320 1325

Ile Glu Leu Lys Asp Arg Lys Arg Pro Ser Glu Leu Arg Arg Ile Thr  
1330 1335 1340

Ser Gln Val Lys Tyr Ala Gly Ser Glu Val Ala Ser Thr Ser Glu Val  
1345 1350 1355 1360

Ile Ile Asp Ile Ala Ile  
1365 1370 1375 1380 1385 1390 1395 1400 1405 1410 1415 1420 1425 1430

Ala Ile Arg Ile Ala Ile  
1435 1440 1445 1450 1455 1460 1465 1470 1475 1480 1485 1490 1495 1500

Leu Ala Arg Asn Leu Val Arg Tyr Val Gln Gly Leu Lys Lys Lys Lys  
1495 1500 1505 1510 1515 1520 1525 1530 1535 1540 1545 1550 1555 1560

Val Ile Val Ile Arg Val Ile  
1565 1570 1575 1580 1585 1590 1595 1600 1605 1610 1615 1620 1625 1630 1635

Leu Cys Asp Leu Ala Pro Glu Ala Pro Ala Pro Thr Gln His Pro Pro  
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Met Ala Gln Val Thr Val Gly Ser Glu Leu Leu Gly Val Ser Ser Pro  
1475 1480 1485

Gly Pro Lys Arg Asn Ser Met Val Leu Asp Val Val Phe Val Leu Glu  
1490 1495 1500

Gly Ser Asp Lys Ile Gly Glu Ala Asn Phe Asn Lys Ser Arg Glu Phe  
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Met Glu Glu Val Ile Gln Arg Met Asp Val Gly Gln Asp Arg Ile His  
1525 1530 1535

Val Thr Val Leu Gln Tyr Ser Tyr Met Val Thr Val Glu Tyr Thr Phe  
1540 1545 1550

Ser Glu Ala Gln Ser Lys Gly Glu Val Leu Gln Gln Val Arg Asp Ile  
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Arg Tyr Arg Gly Gly Asn Arg Thr Asn Thr Gly Leu Ala Leu Gln Tyr  
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Ile Ser Glu His Ser Phe Ser Val Ser Gln Gly Asp Arg Glu Gln Val  
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Pro Asn Leu Val Tyr Met Val Thr Gly Asn Pro Ala Ser Asp Glu Ile  
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Lys Arg Met Pro Gly Asp Ile Gln Val Val Pro Ile Gly Val Gly Pro  
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His Ala Asn Val Gln Glu Leu Glu Lys Ile Gly Trp Pro Asn Ala Pro  
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Ile Leu Ile His Asp Phe Glu Met Leu Pro Arg Glu Ala Pro Asp Leu  
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Asp Thr Ile Asp Thr  
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Asp Thr Thr Ile Asp Thr  
1685 1690 1695 1700

Ile Asp Thr  
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Met Gln Gln Glu Gly Gly Pro Ser Glu Ile Gly Asp Ala Leu Ser Phe  
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Ala Val Arg Tyr Val Thr Ser Glu Val His Gly Ala Arg Pro Gly Ala  
1780 1785 1790

Ser Lys Ala Val Val Ile Leu Val Thr Asp Val Ser Val Asp Ser Val  
1795 1800 1805

Asp Ala Ala Ala Glu Ala Ala Arg Ser Asn Arg Val Thr Val Phe Pro  
1810 1815 1820

Ile Gly Ile Gly Asp Arg Tyr Ser Glu Ala Gln Leu Ser Ser Leu Ala  
1825 1830 1835 1840

Gly Pro Lys Ala Gly Ser Asn Met Val Arg Leu Gln Arg Ile Glu Asp  
1845 1850 1855

Leu Pro Thr Val Ala Thr Leu Gly Asn Ser Phe Phe His Lys Leu Cys  
1860 1865 1870

Ser Gly Phe Asp Arg Val Cys Val Asp Glu Asp Gly Asn Glu Lys Arg  
1875 1880 1885

Pro Gly Asp Val Trp Thr Leu Pro Asp Gln Cys His Thr Val Thr Cys  
1890 1895 1900

Leu Pro Asp Gly Gln Thr Leu Leu Lys Ser His Arg Val Asn Cys Asp  
1905 1910 1915 1920

Arg Gly Pro Arg Pro Ser Cys Pro Asn Gly Gln Pro Pro Leu Arg Val  
1925 1930 1935

Glu Glu Thr Cys Gly Cys Arg Trp Thr Cys Pro Cys Val Cys Met Gly  
1940 1945 1950

Ser Ser Thr Arg His Ile Val Thr Phe Asp Gly Gln Asn Phe Lys Leu  
1955 1960 1965

Thr Asp Asp Tyr Asp Lys Val Ile Ile Ala Asn Asp Asp Lys Asp Ile  
1970 1975 1980

Gly Val Ile Ile Ile Asp Gly Ala Asp Asp Phe Asp Asp Ala Ile Ile Ile  
1985 1990 1995 2000

Cys Met Lys Ser Ile Glu Val Lys His Asp Gly Leu Ser Val Gln Leu  
2005 2010 2015 2020

Asp Ser Asp Asp Met Lys Met Thr Val Asn Asp Asp Leu Val Val Asp Phe  
2025 2030 2035 2040

Asn Asn Glu Phe Gln Leu Gln Leu Ser Pro Arg Thr Phe Ala Ser Lys  
2065 2070 2075 2080

Thr Tyr Gly Leu Cys Gly Ile Cys Asp Glu Asn Gly Ala Asn Asp Phe  
2085 2090 2095

Ile Leu Arg Asp Gly Thr Val Thr Asp Trp Lys Ala Leu Ile Gln  
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Glu Trp Thr Val Gln Gln Leu Gly Lys Thr Ser Gln Pro Val His Glu  
2115 2120 2125

Glu Gln Cys Pro Val Ser Glu Phe Phe His Cys Gln Val Leu Leu Ser  
2130 2135 2140

Glu Leu Phe Ala Glu Cys His Lys Val Leu Ala Pro Ala Thr Phe Tyr  
2145 2150 2155 2160

Ala Met Cys Gln Pro Asp Ser Cys His Pro Lys Lys Val Cys Glu Ala  
2165 2170 2175

Ile Ala Leu Tyr Ala His Leu Cys Arg Thr Lys Gly Val Cys Val Asp  
2180 2185 2190

Trp Arg Arg Ala Asn Phe Cys Ala Met Ser Cys Pro Pro Ser Leu Val  
2195 2200 2205

Tyr Asn His Cys Glu His Gly Cys Pro Arg Leu Cys Glu Gly Asn Thr  
2210 2215 2220

Ser Ser Cys Gly Asp Gln Pro Ser Glu Gly Cys Phe Cys Pro Pro Asn  
2225 2230 2235 2240

Gln Val Met Leu Glu Gly Ser Cys Val Pro Glu Glu Ala Cys Thr Gln  
2245 2250 2255

Cys Ile Ser Glu Asp Gly Val Arg His Gln Phe Leu Glu Thr Trp Val  
2260 2265 2270

Ile Ala Arg Ile Ile Tyr Ile Ile Tyr Thr Tyr Ile Ile Ile Ile Ile Ile  
2275 2280 2285 2290 2295 2300

Val Asn Tyr Thr Ile Ile Ile Ile Ile Tyr Ala Ile Ile Ile Ile Ile Ile Ile  
2305 2310 2315 2320 2325 2330

Tyr Pro Cys Glu Val Ala Arg Leu Arg; Gln Asn Ala Val Gln Tyr Lys  
2335 2340 2345 2350 2355 2360

Ile Ile Tyr Ile Ile Tyr Val Val Asp Leu Val Ser Tyr Asn Ile Ile Ile  
2365 2370 2375 2380 2385 2390

Arg Glu Ser Pro Pro Ser Cys Pro Pro His Arg Thr Pro Ala Leu Arg  
2370 2375 2380

Lys Thr Gln Cys Cys Asp Glu Tyr Glu Cys Ala Cys Asn Cys Val Asn  
2385 2390 2395 2400

Ser Thr Val Ser Cys Pro Leu Gly Tyr Leu Ala Ser Ala Val Thr Asn  
2405 2410 2415

Asp Cys Gly Cys Thr Thr Thr Cys Phe Pro Asp Lys Val Cys Val  
2420 2425 2430

His Arg Gly Thr Ile Tyr Pro Val Gly Gln Phe Trp Glu Glu Ala Cys  
2435 2440 2445

Asp Val Cys Thr Cys Thr Asp Leu Glu Asp Ser Val Met Gly Leu Arg  
2450 2455 2460

Val Ala Gln Cys Ser Gln Lys Pro Cys Gln Asp Asn Cys Leu Ser Gly  
2465 2470 2475 2480

Phe Thr Tyr Val Leu His Glu Gly Glu Cys Cys Gly Arg Cys Leu Pro  
2485 2490 2495

Ser Ala Cys Glu Val Val Thr Gly Ser Pro Arg Gly Asp Ala Gln Ser  
2500 2505 2510

His Trp Lys Asn Val Gly Ser His Trp Ala Ser Pro Asp Asn Pro Cys  
2515 2520 2525

Leu Ile Asn Glu Cys Val Arg Val Lys Glu Glu Val Phe Val Gln Gln  
2530 2535 2540

Arg Asn Val Ser Cys Pro Gln Leu Asn Val Pro Thr Cys Pro Thr Gly  
2545 2550 2555 2560

Phe Gln Leu Ser Cys Lys Thr Ser Glu Cys Cys Pro Thr Cys His Cys  
2565 2570 2575

Leu Ile Asn Asn Asp Ile Asp Val Asp Thr Thr Tyr Asp Tyr Thr Val Ile  
2580 2585 2590 2595

Val Gly Val Ile Ser Gly Phe Ile Val Glu Ile Asp Ile Asp Thr Thr Tyr  
2595 2600 2605 2610

Asn Ala Asp Ile Ile Asp Ile Asp Tyr Tyr Asp Ile Asp Asn Asp Ile Asp  
2615 2620 2625 2630

|   |      |      |      |
|---|------|------|------|
| Ser His Phe Cys Lys Val Asn Glu Arg Gly Glu Tyr Ile Thr Glu Lys |      |      |      |
| 2675  | 2680 | 2685 |      |
| Arg Val Thr Gly Cys Pro Pro Phe Asp Glu His Lys Cys Leu Ala Glu |      |      |      |
| 2690  | 2695 | 2700 |      |
| Gly Gly Lys Ile Met Lys Ile Pro Gly Thr Cys Cys Asp Thr Cys Glu |      |      |      |
| 2705  | 2710 | 2715 | 2720 |
| Glu Pro Glu Cys Lys Asp Ile Ile Ala Lys Leu Gln Arg Val Lys Val |      |      |      |
| 2725  | 2730 | 2735 |      |
| Gly Asp Cys Lys Ser Glu Glu Glu Val Asp Ile His Tyr Cys Glu Gly |      |      |      |
| 2740  | 2745 | 2750 |      |
| Lys Cys Ala Ser Lys Ala Val Tyr Ser Ile His Met Glu Asp Val Gln |      |      |      |
| 2755  | 2760 | 2765 |      |
| Asp Gln Cys Ser Cys Cys Ser Pro Thr Gln Thr Glu Pro Met Gln Val |      |      |      |
| 2770  | 2775 | 2780 |      |
| Ala Leu Arg Cys Thr Asn Gly Ser Leu Ile Tyr His Glu Ile Leu Asn |      |      |      |
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| Ala Ile Glu Cys Arg Cys Ser Pro Arg Lys Cys Ser Lys             |      |      |      |
| 2805  | 2810 |      |      |

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• 111-63



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521.2 14  
521.3 15  
521.4 16  
521.5 17  
521.6 18  
521.7 19  
521.8 20  
521.9 21

1911-1912  
1911-1912  
1911-1912  
1911-1912

|  |    |  |
|--|----|--|
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| 10311> Cysaa   | 15 |  |
| 401> 20  |    |  |
| 211> 16  |    |  |
| 212> DNA   |    |  |
| 213> Canis familiaris  |    |  |
| 402> 20  |    |  |
| 10311> tsqaa   | 16 |  |
| 210> 21  |    |  |
| 211> 218   |    |  |
| 212> DNA   |    |  |
| 213> Canis familiaris  |    |  |
| 403> 21  |    |  |
| gtccctgtggg agcagtgcata gctccatgaag agtgcctcggt tgtttgcggc ctgcaccccg 60 |    |  |
| ctgggtggacc ctgagacttt tgccgcctgt tggaaaggaa ctgtgtgcac ctgtgtccag 120   |    |  |
| ggatggagt gcccattgtgc ggtctctgtg gagtaacccc gggctgtgc ccaacaggaa 180     |    |  |
| atgtgtgtgt acggatggac cggccacagg gtctgtgg 218                            |    |  |
| 404> 22  |    |  |
| 211> 73  |    |  |
| 212> PRT   |    |  |
| 213> Canis familiaris  |    |  |
| 405> 22  |    |  |
| Val Leu Trp Glu Gln Cys Gln Leu Leu Lys Ser Ala Ser Val Phe Ala          |    |  |
| 1 5 10 15  |    |  |
| Arg Cys His Pro Leu Val Asp Pro Glu Ile Phe Val Ala Leu Cys Glu          |    |  |
| 20 25 30 35  |    |  |
| Arg Thr Leu Cys Thr Cys Val Gln Gly Met Ile Cys Pro Cys Ala Val          |    |  |
| 35 40 45   |    |  |
| Leu Leu Glu Tyr Ala Arg Ala Cys Ala Gln Gln Gly Ile Val Leu Tyr          |    |  |
| 50 55 60   |    |  |
| Arg Ile Ile Asp Glu Asp Val Val Asp                                      |    |  |
| 65 70 75   |    |  |
| 214> DNA   |    |  |
| 215> Canis familiaris  |    |  |

